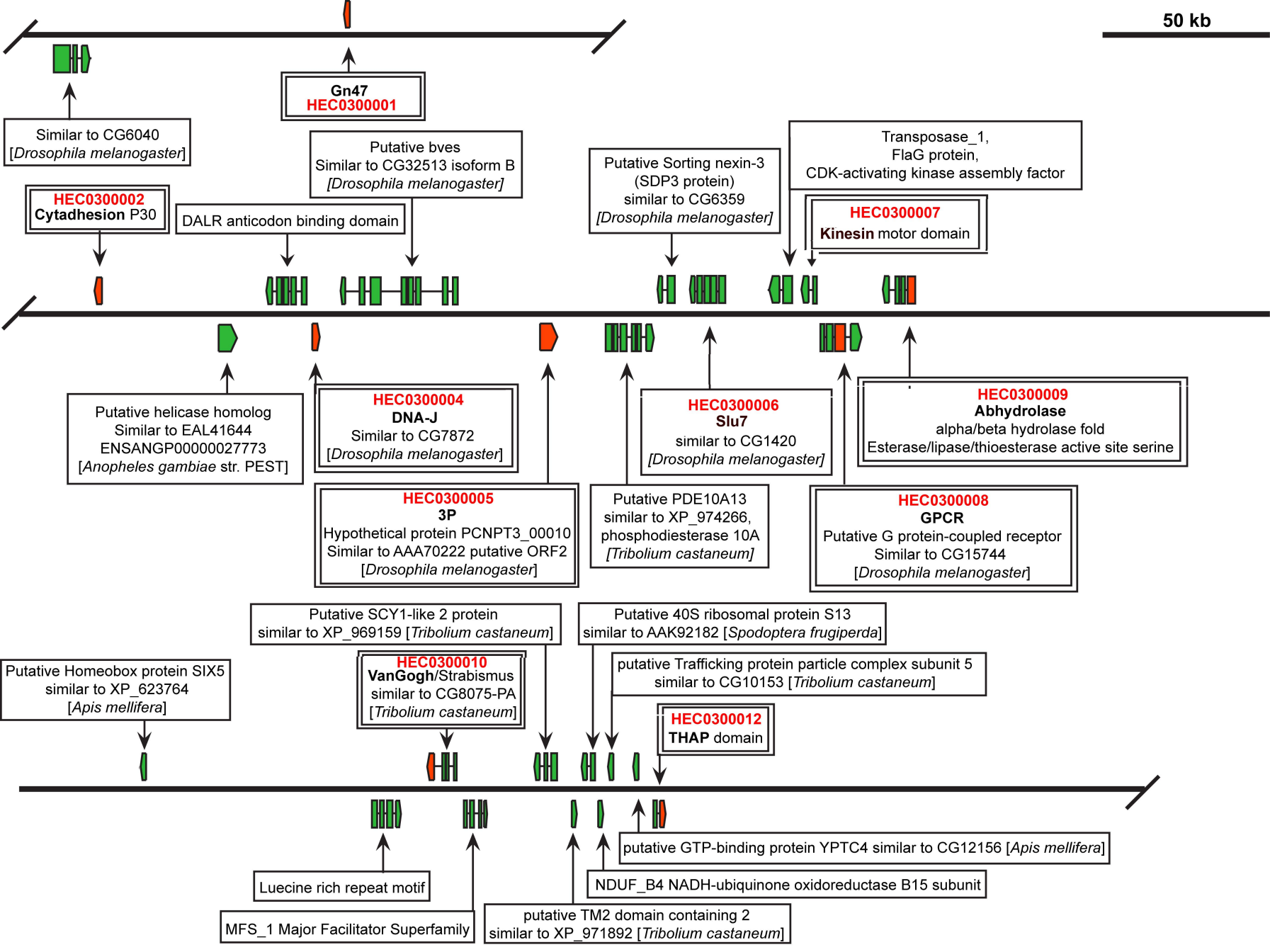
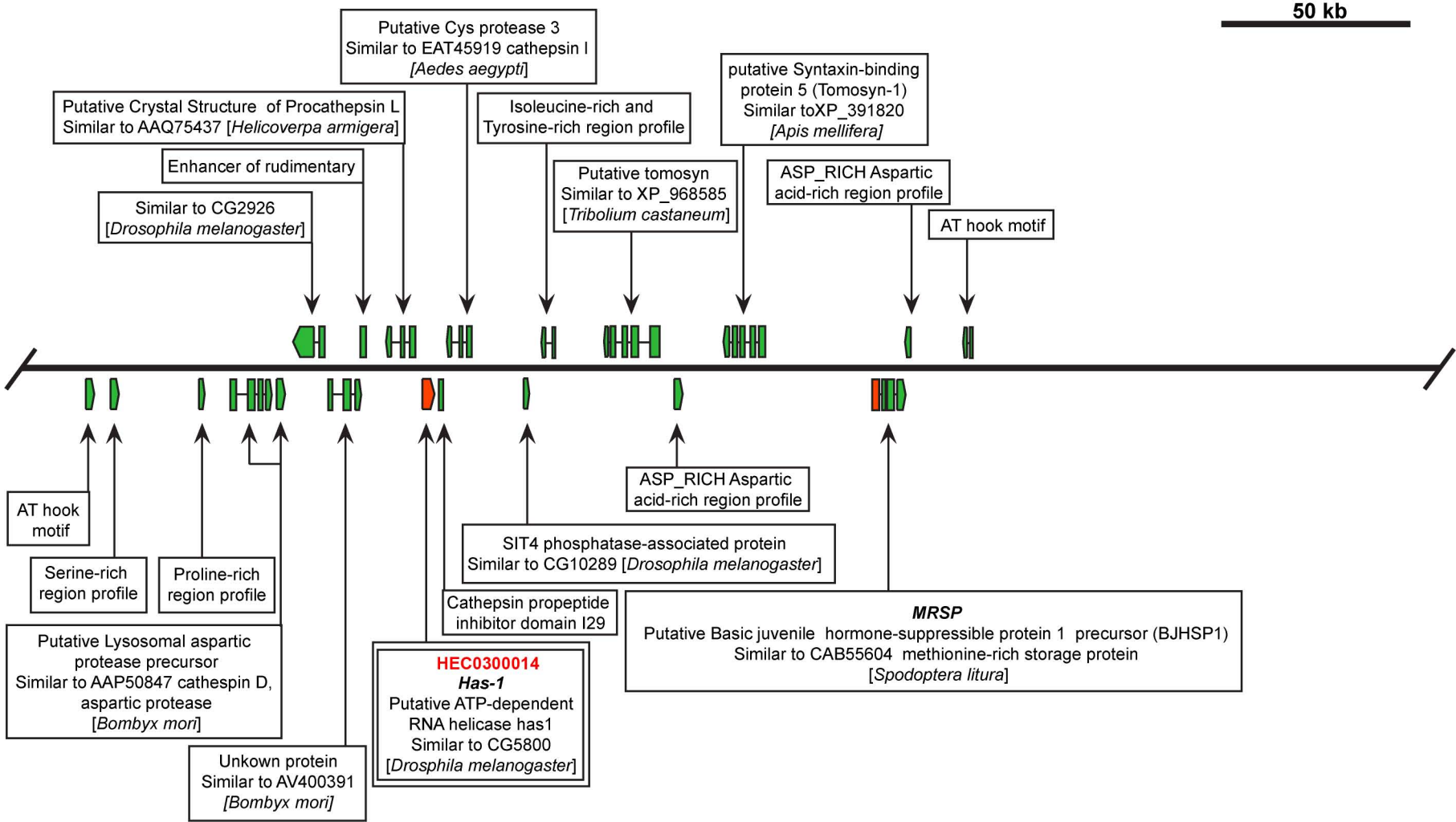
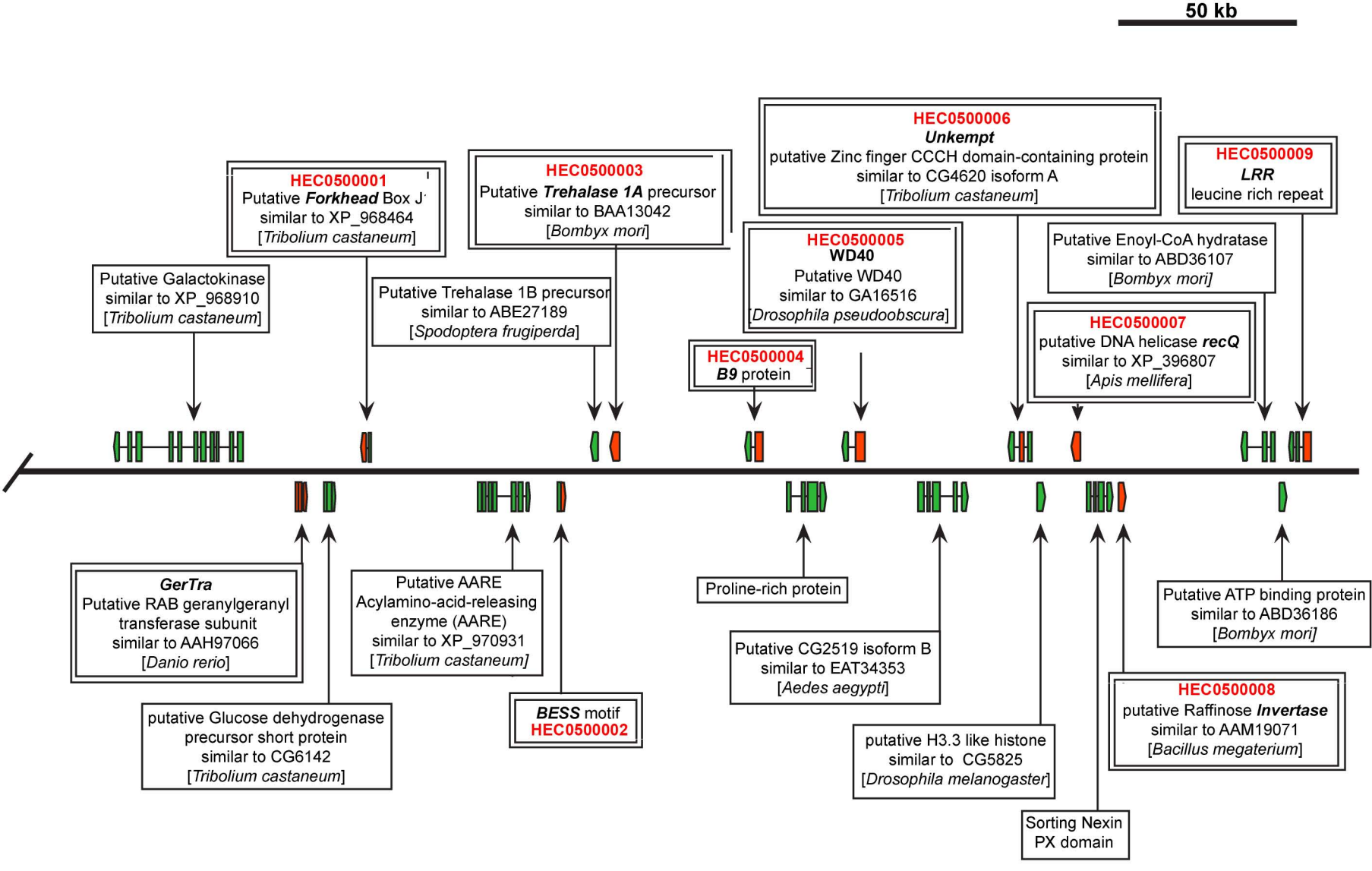

50 kb
Gn47
HEC0300001
Similar to CG6040
[Drosophila melanogaster]
Putative bves
Similar to CG32513 isoform B
[Drosophila melanogaster]
HEC0300002
Cytadhesion P30
DALR anticodon binding domain
Putative Sorting nexin-3
(SDP3 protein)
similar to CG6359
[Drosophila melanogaster]
Transposase_1,
FlaG protein,
CDK-activating kinase assembly factor
HEC0300007
Kinesin motor domain
Putative helicase homolog
Similar to EAL41644
ENSANGP00000027773
[Anopheles gambiae str. PEST]
HEC0300004
DNA-J
Similar to CG7872
[Drosophila melanogaster]
HEC0300006
Slu7
similar to CG1420
[Drosophila melanogaster]
HEC0300009
Abhydrolase
alpha/beta hydrolase fold
Esterase/lipase/thioesterase active site serine
HEC0300005
3P
Hypothetical protein PCNPT3_00010
Similar to AAA70222 putative ORF2
[Drosophila melanogaster]
Putative PDE10A13
similar to XP_974266,
phosphodiesterase 10A
[Tribolium castaneum]
HEC0300008
GPCR
Putative G protein-coupled receptor
Similar to CG15744
[Drosophila melanogaster]
Putative SCY1-like 2 protein
similar to XP_969159 [Tribolium castaneum]
Putative 40S ribosomal protein S13
similar to AAK92182 [Spodoptera frugiperda]
Putative Homeobox protein SIX5
similar to XP_623764
[Apis mellifera]
HEC0300010
VanGogh/Strabismus
similar to CG8075-PA
[Tribolium castaneum]
putative Trafficking protein particle complex subunit 5
similar to CG10153 [Tribolium castaneum]
HEC0300012
THAP domain
putative GTP-binding protein YPTC4 similar to CG12156 [Apis mellifera]
Luecine rich repeat motif
NDUF_B4 NADH-ubiquinone oxidoreductase B15 subunit
MFS_1 Major Facilitator Superfamily
putative TM2 domain containing 2
similar to XP_971892 [Tribolium castaneum]


50 kb
Putative Cys protease 3
Similar to EAT45919 cathepsin l
[Aedes aegypti]
Putative Crystal Structure of Procathepsin L
Similar to AAQ75437 [Helicoverpa armigera]
Isoleucine-rich and
Tyrosine-rich region profile
putative Syntaxin-binding
protein 5 (Tomosyn-1)
Similar toXP_391820
[Apis mellifera]
Enhancer of rudimentary
Putative tomosyn
Similar to XP_968585
[Tribolium castaneum]
ASP_RICH Aspartic
acid-rich region profile
Similar to CG2926
[Drosophila melanogaster]
AT hook motif
AT hook
motif
Serine-rich
region profile
Proline-rich
region profile
ASP_RICH Aspartic
acid-rich region profile
SIT4 phosphatase-associated protein
Similar to CG10289 [Drosophila melanogaster]
Cathepsin propeptide
inhibitor domain I29
MRSP
Putative Basic juvenile hormone-suppressible protein 1 precursor (BJHSP1)
Similar to CAB55604 methionine-rich storage protein
[Spodoptera litura]
Putative Lysosomal aspartic
protease precursor
Similar to AAP50847 cathespin D,
aspartic protease
[Bombyx mori]
HEC0300014
Has-1
Putative ATP-dependent
RNA helicase has1
Similar to CG5800
[Drosphila melanogaster]
Unkown protein
Similar to AV400391
[Bombyx mori]


50 kb
HEC0500006
Unkempt
putative Zinc finger CCCH domain-containing protein
similar to CG4620 isoform A
[Tribolium castaneum]
HEC0500009
LRR
leucine rich repeat
HEC0500001
Putative Forkhead Box J
similar to XP_968464
[Tribolium castaneum]
HEC0500003
Putative Trehalase 1A precursor
similar to BAA13042
[Bombyx mori]
HEC0500005
WD40
Putative WD40
similar to GA16516
[Drosophila pseudoobscura]
Putative Enoyl-CoA hydratase
similar to ABD36107
[Bombyx mori]
Putative Galactokinase
similar to XP_968910
[Tribolium castaneum]
Putative Trehalase 1B precursor
similar to ABE27189
[Spodoptera frugiperda]
HEC0500007
putative DNA helicase recQ
similar to XP_396807
[Apis mellifera]
HEC0500004
B9 protein
GerTra
Putative RAB geranylgeranyl
transferase subunit
similar to AAH97066
[Danio rerio]
Putative AARE
Acylamino-acid-releasing
enzyme (AARE)
similar to XP_970931
[Tribolium castaneum]
Proline-rich protein
Putative ATP binding protein
similar to ABD36186
[Bombyx mori]
Putative CG2519 isoform B
similar to EAT34353
[Aedes aegypti]
HEC0500008
putative Raffinose Invertase
similar to AAM19071
[Bacillus megaterium]
putative Glucose dehydrogenase
precursor short protein
similar to CG6142
[Tribolium castaneum]
BESS motif
HEC0500002
putative H3.3 like histone
similar to CG5825
[Drosophila melanogaster]
Sorting Nexin
PX domain